



GGC	ACG	AGG	GGC	GGC	CTG	CGG	CGC	AGAG	CGG	GAG	ATG	CAG	CGG	CTT	GGG	GCC	ACC	CTG	CTG	TGC	10
L	L	L	A	A	V	P	T	A	P	A	P	A	P	A	P	T	A	T	S	A	30
CTG	CTG	CTG	GCG	GCG	GTC	CCC	ACG	GCC	CCC	GCG	CCC	GCG	CCC	GCT	CCG	ACG	GCG	ACC	TCG	GCT	127
P	V	K	P	G	P	A	L	S	Y	P	Q	E	E	A	T	L	N	E	M	50	
CCA	GTC	AAG	CCC	GGC	CCG	GCT	CTC	AGC	TAC	CCG	CAG	GAG	GAG	GCC	ACC	CTC	AAT	GAG	ATG	187	
F	R	E	V	E	E	L	M	E	D	T	Q	H	K	L	R	S	A	V	E	70	
TTT	CGC	GAG	GTT	GAG	GAA	CTG	ATG	GAG	GAC	ACG	CAG	CAC	AAA	TTG	CGC	AGC	GCG	GTG	GAA	247	
E	M	E	A	E	E	A	A	A	K	A	S	S	E	V	N	L	A	N	L	90	
GAG	ATG	GAG	GCA	GAA	GAA	GCT	GCT	GCT	AAA	GCA	TCA	TCA	GAA	GTG	AAC	CTG	GCA	AAC	TTA	307	
P	P	S	Y	H	N	E	T	N	T	D	T	K	V	G	N	N	T	I	H	110	
CCCT	CCC	AGC	TAT	CAC	AAT	GAG	ACC	AAC	ACA	GAC	ACG	AAG	GTT	GGA	AAT	AAT	ACC	ATC	CAT	367	
V	H	R	E	I	H	K	I	T	N	N	Q	T	G	Q	M	V	F	S	E	130	
GTG	CAC	CGA	GAA	ATT	CAC	AAG	ATA	ACC	AAC	AAC	CAG	ACT	GGA	CAA	ATG	GTC	TTT	TCA	GAG	427	
T	V	I	T	S	V	G	D	E	E	G	R	R	S	H	E	C	I	I	D	150	
ACA	GTT	ATC	ACA	TCT	GTG	GGA	GAC	GAA	GAA	GGC	AGA	AGG	AGC	CAC	GAG	TGC	ATC	ATC	GAC	487	
E	D	C	G	P	S	M	Y	C	Q	F	A	S	F	Q	Y	T	C	Q	P	170	
GAG	GAC	TGT	GGG	CCC	AGC	ATG	TAC	TGC	CAG	TTT	GCC	AGC	TTC	CAG	TAC	ACC	TGC	CAG	CCA	547	
C	R	G	Q	R	M	L	C	T	R	D	S	E	C	C	G	D	Q	L	C	190	
TGC	CGG	GGC	CAG	AGG	ATG	CTC	TGC	ACC	CGG	GAC	AGT	GAG	TGC	TGT	GGA	GAC	CAG	CTG	TGT	607	

[illegible]

FIG. 1C

ATCTGGACCAGGCTGTGGGTAGATGTGCAATAGAAAATAGCTAATTATTTCCTCCCANGTGTGTGCTTTAAAGCGTGGGCTG 1169
ACCAGGCTTCTTCCTACATCTTCTTCCAGTAAGTTTCCCTCTGGCTTGACAGCATGAGGTGTGTGTCATTTGTTTCAG 1248
CTCCCCCAGGCTGTTCTCCAGGCTTCACAGTCTGGTGCTTGGGAGAGTCAGGCAGGGTTAAACTGCAGGAGCAGTTTGC 1327
CACCCCTGTCCAGATTATTGGCTGCTTTGGCTCTACCAAGTTGGCAGACAGCCGTTTGTTCACATGGCTTTTGATAATTG 1406
TTTGAGGGGAGGAGATGGAAACAATGTGGAGTCTCCCTCTGATTGGTTTGGGGAATGTGGAGAAAGAGTGCCCTGCTT 1485
TGCAAAACATCAACCTGGCAAAAATGCAACAAATGAATTTTCCACGCAGTTCTTTCCATGGGCATAGGTAAGCTGTGCCT 1564
TCAGCTGTTGCAGATGAATGTCTGTTCACCCCTGCATTACATGTGTTTATTTCATCCAGCAGTGTGTGCTCAGCTCCTAC 1643
CTCTGTGCCAGGCGAGCATTTTCATATCCAAGATCAATTCCCTCTCTCAGCACAGCCTGGGGAGGGGTTCATTGTTCTC 1722
CTCGTCCATCAGGGATTTCAGAGGCTCAGAGACTGCAAGCTGCTTGCCCAAGTCACACAGCTAGTGAAGACCAGAGCAG 1801
TTTTCATCTGGTGTGACTCTAAGCTCAGTGTCTCTCCACTACCCACACAGCCTTGGTGCCACCAAAAGTGCTCCCC 1880
AAAAGGAAGGAGAATGGGATTTTCTTTTGAGGCATGCACATCTGGAATTAAGGTCAAACTAATTCTCACATCCCTCTA 1959
AAAGTAAACTACTGTTAGGAACAGCAGTGTCTCACAGTGTGGGGCAGCCGTCCTTCTAATGAAGACAATGATATTGAC 2038
ACTGTCCCCTCTTTGGCAGTTGCATTAGTAACTTTGAAAGGTATATGACTGAGCGTAGCATACAGGTTAACCTGCAGAAA 2117
CAGTACTTAGGTAATTGTAGGGCGAGGATTATAAATGAATTTTGCAAAATCACTTAGCAGCAACTGAAGACAATTATCA 2196

FIG. 1D

ACCACGTGGAGAAAATCAAACCGAGCAGGGCTGTGTGAAACATGGTTGTAATATGCCGACTGCCGAACACTGAACCTCTACG 2275

CCACTCCACAAATGATGTTTTTCAGGGTGTCAATGGACTGTGGCCACCATGTATTTCATCCAGAGTTCCTTAAAGTTTAAAGTT 2354

GCACATGATTGTATAAGCATGCTTTCTTTGAGTTTAAATTATGTATAAACATAAAGTTGCATTTAGAAATCAAGCATAA 2433

ATCACTTCAACTGCTAAAAAATAAAAAAATAAAAAAATAAAAAAATAAAAAAATAAAAAAATAAAAAAATAAAAAAATAAAAAA 2479

FIG. 2A

GAATTGGCAGGAGACGACGTGCTGAGCTGCCAGCTTAGTGGAAGCTCTGCTCTGGTGGAGAGCAGCCTCGCTTTG	79
M V A A V L L G	
GTGACGCACAGTCTGGGACCCCTCCAGGAGCCCGGATTGAAGG ATG GTG GCG GCC GTC CTG CTG GGG	8 148
L S W L C S P L G A L V L D F N N I R S	28
CTG AGC TGG CTC TGC TCT CCC CTG GGA GCT CTG GTC CTG GAC TTC AAC AAC ATC AGG AGC	208
S A D L H G A R K G S Q C L S D T D C N	48
TCT GCT GAC CTG CAT GGG GCC CGG AAG GGC TCA CAG TGC CTG TCT GAC ACG GAC TGC AAT	268
T R K F C L Q P R D E K P F C A T C R G	68
ACC AGA AAG TTC TGC CTC CAG CCC CGC GAT GAG AAG CCG TTC TGT GCT ACA TGT CGT GGG	328
L R R R C Q R D A M C C P G T L C V N D	88
TTG CGG AGG AGG TGC CAG CGA GAT GCC ATG TGC TGC CCT GGG ACA CTC TGT GTG AAC GAT	388
V C T T M E D A T P I L E R Q L D E Q D	108
GTT TGT ACT ACG ATG GAA GAT GCA ACC CCA ATA TTA GAA AGG CAG CTT GAT GAG CAA GAT	448
G T H A E G T T G H P V Q E N Q P K R K	128
GGC ACA CAT GCA GAA GGA ACA ACT GGG CAC CCA GTC CAG GAA AAC CAA CCC AAA AGG AAG	508
P S I K K S Q G R K G Q E G E S C L R T	148
CCA AGT ATT AAG AAA TCA CAA GSC AGG AAG GGA CAA GAG GGA GAA AGT TGT CTG AGA ACT	568

FIG. 2B

```
F D C G P G L C C A R H F W T K I C K P 168
TTT GAC TGT GGC CCT GGA CTT TGC TGT GCT CGT CAT TTT TGG ACG AAA ATT TGT AAG CCA 628

V L L E G Q V C S R R G G G CAT AAA GAC ACT GCT CAA Q A P 188
GTC CTT TTG GAG GGA CAG GTC TGC TCC AGA AGA GGG CAT AAA GAC ACT GCT CAA GCT CCA 688

E I F Q R C D C G P G L L C R S Q L T S 208
GAA ATC TTC CAG CGT TGC GAC TGT GGC CCT GGA CTA CTG TGT CGA AGC CAA TTG ACC AGC 748

N R Q H A R L R V C Q K I E K L * 225
AAT CGG CAG CAT GCT CGA TTA AGA GTA TGC CAA AAA ATA GAA AAG CTA TAA 799

ATATTTCAAATAAAGAAGAAATCCACATTGCATAAAAAAAAAAAAAA 848
```


FIG. 3A

CCGGACGCGTGGCGGACACGGTTTCGTGGGGACCCAGGCTTGCAAAGTGACGGTCA	TTTCTCTTTCTTTCTCTCTCTT	79
M M A L G A A G A T R V F V A M		
GAGTCCTTCTGAG ATG ATG GCT CTG GGC GCA GCG GGA GCT ACC CGG GTC TTT GTC GCG ATG		16
V A A A L G G H P L L G V S A T L N S V		
GTA GCG GCG GCT CTC GGC GGC CAC CCT CTG CTG GGA GTG AGC GCC ACC TTG AAC TCG GTT		140
L N S N A I K N L P P P L G G A A G H P		
CTC AAT TCC AAC GCT ATC AAG AAC CTG CCC CCA CCG CTG GGC GGT GCG GGG CAC CCA		36
G S A V S A A P G I L Y P G G N K Y Q T		
GGC TCT GCA GTC AGC GCC GCG CCG GGA ATC CTG TAC CCG GGC GGC AAT AAG TAC CAG ACC		200
I D N Y Q P Y P C A E D E E C G T D E Y		
ATT GAC AAC TAC CAG CCG TAC CCG TGC GCA GAG GAC GAG GAG TGC GGC ACT GAT GAG TAC		96
C A S P T R G G G A G V Q I C L A C R K		
TGC GCT AGT CCC ACC CGC GGA GGC GAC GCA GGC GTG CAA ATC TGT CTC GCC TGC AGG AAG		116
R R K R C M R H A M C C P G N Y C K N G		
CGC CGA AAA CGC TGC ATG CGT CAC GCT ATG TGC TGC CCC GGC AAT TAC TGC AAA AAT GGA		136
I C V S S D Q N H F R G E I E E T I T E		
ATA TGC GTG TCT TCT GAT CAA AAT CAT TTC CGA GGA GAA ATT GAG GAA ACC ATC ACT GAA		156
560		

FIG. 3B

S F G N D H S T L D G Y S R R T T L S S 176
AGC TTT GGT AAT GAT CAT AGC ACC TTG GAT GGG TAT TCC AGA AGA ACC ACC TTG TCT TCA 620

K M Y H T K G Q E G S V C L R S S D C A 196
AAA ATG TAT CAC ACC AAA GGA CAA GAA GGT TCT GTT TGT CTC CGG TCA TCA GAC TGT GCC 680

S G L C C A R H F W S K I C K P V L K E 216
TCA GGA TTG TGT TGT GCT AGA CAC TTC TGG TCC AAG ATC TGT AAA CCT GTC CTG AAA GAA 740

G Q V C T K H R R K G S H G L E I F Q R 236
GGT CAA GTG TGT ACC AAG CAT AGG AGA AAA GGC TCT CAT GGA CTA GAA ATA TTC CAG CGT 800

C Y C G E G L S C R I Q K D H H Q A S N 256
TGT TAC TGT GGA GAA GGT CTG TCT TGC CGG ATA CAG AAA GAT CAC CAT CAA GCC AGT AAT 860

S S R L H T C Q R H * 267
TCT TCT AGG CTT CAC ACT TGT CAG AGA CAC TAA 893

ACCAGCTATCCAAAAATGCAGTGAACTCCTTTTATATAATAGATGCTATGAAAAACCTTTTATGACCTTCATCAACTCAAT 972
CCTAAGGATATACAAGTTCTGTGTTTCAGTTAAGCATTCCAATAACACCTTCCAAAAACCTGGAGTGTAAAGAGCTTG 1051
TTTCTTTATGGAACTCCCTGTGATTGCAGTAAATTAAGTATGTAATTCAGTGTGGCACTTACCTGTAAATGCA 1130
ATGAAACTTTTAAATTAATTTTCTAAAGGTGCTGCACTGCCTATTTTCTCTTGTATGTAATTTTGTACACATTGA 1209
TTGTTATCTTGACTGACAAATATCTATATTGAACCTGAAGTAAATCATTTTCAGCTTATAGTTCTTAAAGCATAAACCCT 1288
TTACCCCATTTNATTTCTAGAGTCNAGAACGCAAGGATCTCTTGAATGACAAATGATAGGTACCTAAATGTAAACATGA 1367
AAATACTAGCTTATTTTCTGAAATGTACTATCTTAATGCTTAAATTAATTTCCCTTTAGGCTGTGATAGTTTGTGAAA 1446
TAAAAATTAAACATTTAATATCATGAAATGKTTAAGTAGACATAAAAAAAGGGGGGGGGCTAGTA 1525
CTAG 1529

FIG. 4

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E F G T R V G R Y C H S P H Q G S S A C 20
GAA TTC GGC ACG AGG GTT GGG AGG TAT TGC CAC AGT CCC CAC CAA GGA TCA TCG GCC TGC 60

M V C R R K K K R C H R D G M C C P S T 40
ATG GTG TGT CGG AGA AAA AAG AAG CGC TGC CAC CGA GAT GGC ATG TGC TGC CCC AGT ACC 120

R C N N G I C I P V T E S I L T P H I P 60
CGC TGC AAT AAT GGC ATC TGT ATC CCA GTT ACT GAA AGC ATC TTA ACC CCT CAC ATC CCG 180

A L D G T R H R D R N H G H Y S N H D L 80
GCT CTG GAT GGT ACT CGG CAC AGA GAT CGA AAC CAC GGT CAT TAC TCA AAC CAT GAC TTG 240

G W Q N L G R P H T K M S H I K G H E G 100
GGA TGG CAG AAT CTA GGA AGA CCA CAC ACT AAG ATG TCA CAT ATA AAA GGG CAT GAA GGA 300

D P C L R S S D C I E G F C C A R H F W 120
GAC CCC TGC CTA CGA TCA TCA GAC TGC ATT GAA GGG TTT TGC TGT GCT CGT CAT TTC TGG 360

T K I C K P V L H Q G E V C T K Q R K K 140
ACC AAA ATC TGC AAA CCA GTG CTC CAT CAG GGG GAA GTC TGT ACC AAA CAA CGC AAG AAG 420

G S H G L E I F Q R C D C A K G L S C K 160
GGT TCT CAT GGG CTG GAA ATT TTC CAG CGT TGC GAC TGT GCG AAG GGC CTG TCT TGC AAA 480

V W K D A T Y S S K A R L H V C Q K I * 180
GTA TGG AAA GAT GCC ACC TAC TCC TCC AAA GCC AGA CTC CAT GTG TGT CAG AAA ATT TGA 540

TCACCATTGAGGAACATCATCAATTGCAGACTGTGAAGTTGTGTATTTAATGCATTATAGCATGTGGAAAAATAAGGTT 619
CAGATGCAGAAAGATGGCTAAAAATAAGAAACGTGATAAGAAATATAGATGATCACAAAAAAAAGATGCGG 698
CCGC 702
```


FIG. 5A

CTCGAGGCCAAATTCGGCACGAGCGCGGTGTGGTCTAGCATAAAGCGGAGCCAGAAAGGGCGGGGT	ATG	M	1
			77
G E A S P P A P A R R H L L V L L L L L	L		21
GGA GAA GCC TCC CCA CCT GCC CCC GCA AGG CGG CAT CTG CTG GTC CTG CTC CTC	CTC		137
S T L V I P S A A A P I H D A D A Q E S			41
TCT ACC CTG GTG ATC CCC TCC GCT GCA GCT CCT ATC CAT GAT GCT GAC GCC CAA GAG AGC	AGC		197
S L G L T G L Q S L L L Q G F S R L F L K			61
TCC TTG GGT CTC ACA GGC CTC CAG AGC CTA CTC CAA GGC TTC AGC CGA CTT TTC CTG AAA	AAA		257
G N L L R G I D S L F S A P M D F R G L	L		81
GGT AAC CTG CTT CGG GGC ATA GAC AGC TTA TTC TCT GCT GAC TTC CGG GGC CTC	CTC		317
P G N Y H K E E N Q E H Q L G N N T L S	S		101
CCT GGG AAC TAC CAC AAA GAG AAC CAG GAG CAC CAG CTG GGG AAC AAC ACC CTC TCC	TCC		377
S H L Q I D K M T D N K T G E V L I S E	E		121
AGC CAC CTC CAG ATC GAC AAG ATG ACC GAC AAC AAG ACA GGA GAG GTG CTG ATC TCC GAG	GAG		437
N V V A S I Q P A E G S F E G D L K V P	P		141
AAT GTG GTG GCA TCC ATT CAA CCA GCG GAG GGG AGC TTC GAG GGT GAT TTG AAG GTA CCC	CCC		497
R M E E K E A L V P I Q K A T D S F H T	T		161
AGG ATG GAG GAG AAG GAG GCC CTG GTA CCC ATC CAG AAG GCC ACG GAC AGC TTC CAC ACA	ACA		557
E L H P R V A F W I I K L P R R S H Q	Q		181
GAA CTC CAT CCC CGG GTG GCC TTC TGG ATC ATT AAG CTG CCA CGG AGG TCC CAC CAG	CAG		617

FIG. 5B

D A L E G G G H W L S E K R H R L Q A I R	201
GAT GCC CTG GAG GGC GGC CAC TGG CTC AGC GAG AAG CGA CAC CGC CTG CAG GCC ATC CGG	677
D G L R R K G T H K D V L E E G T E S S	221
GAT GGA CTC CGC AAG GGC ACC CAC AAG GAC GTC CTA GAA GAG GGG ACC GAG AGC TCC TCC	737
H S R L S P R K T H L L Y I L R P S R Q	241
CAC TCC AGG CTG TCC CCC CGA AAG ACC CAC TTA CTG TAC ATC CTC AGG CCC TCT CGG CAG	797
L *	243
CTG TAG	803
GGGTGGGACCGGGGAGCACCTGCCTGTAGCCCCCATCAGACCCCTGCCCAAGCACCATATGGAATAAAGTTCTTCT	882
TACATCTAAAAAATAAAAAAATAAAAAAATAATGGCGGCGCCG	928

FIG. 6A

crsp-2h		
crsp-3h		
crsp-4h		
tango59	MQRLGATLLCLLLAAAVPTAPAPAPTATSAPVKPGPALS		40
Consensus			
crsp-2h		
crsp-3hMMALGAAGATRVFVAMVA		
crsp-4h		
tango59	PQEEATLNEMFREVEELMEDTQHKLRSAVEEMEAEAAAK		80
Consensus		a a	
crsp-2hMVAAVLLGLSWLCS		
crsp-3h	AALGGHPLLGVSATLNSVLNSNAIKNLPPPLGGAAGHPGS		
crsp-4h		
tango59	ASSEVNLANLPPSYHNENTNTDTNVGNNTIHVHREIHKITN		120
Consensus		a n n v s	
crsp-2h	PLGALVLDENNIRSADLHGARKGSQCLSDTDCNTRKFCL		
crsp-3h	AVSAAPGILYPGGNKYQTIDNYQYPYCAEDEECGTDEYCA		
crsp-4hEFGTRVGRYCH		
tango59	NQTGQMVFSETVITSVGDEEGRRSHECIIIDEDCGPSMYCQ		160
Consensus		a s r C DedCgt YC	
crsp-2h	QPRDEKPF...CATCRGLRRRCORDAMCCPGTLCVNDVC		
crsp-3h	SPTRGGDAGVQICLACRKRKRRCMRHAMCCPGNYCKNGIC		
crsp-4h	SPHQGSSA...CMVCRKKRKRCHRDGMCCPSTRCNGIC		
tango59	FASFQYT...CQPCRQRMCLCTRDSECCGDQLCVWGHG		200
Consensus		sP g a C CRg RkRC RDaMCCPgtlCvNGiC	
crsp-2h	TMEDATPILERQLDEQDGTAEHTTGHPVQENQ...PKR		
crsp-3h	VSSDQN..HFRGETIXETI.TESFGNDHSTLD...GYSRR		
crsp-4h	IPVTES..ILTPHIPALDGTNRHRDRNHGHYSNHDLGWQNL		
tango59	TKM.....		240
Consensus		t m il i e dgT g h g r	
crsp-2h	KPSIKKSQGRKGQEGESCLRTEDCGPGLCCARHFWTK..I		
crsp-3h	TTLSSKMYHTKGQEGSVCLRSSDCASGLCCARHFWSK..I		
crsp-4h	GRPHTKMSHIKGEHGDPCCLRSSDCIEGFCCARHFWTK..I		
tango59ATRGSNGTICDNQRDCQPGLCCAFQRGLLFPV		280
Consensus		Km htKGqEG CLRssDC pGLCCARHFWtK I	

FIG. 6B

[illegible]

FIG. 8A

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FGTCGACCCACGGCTCCGCTGTGGCAGCCAGCTACCGTCTGTCGACCATCCAGCTTGACGCTCAGCTTTGTTCATTC 79

      M   Q   R   L   G   G   I   L   L   C   T   L
GAATTGGCGCGCGCCAGCGCGGAACAAAC ATG CAG CGG CTC GGG GGT ATT TTG CTG TGT ACA CTG 12
                                           145

L   A   A   A   V   P   T   A   P   A   P   S   P   T   V   T   W   T   P   A   32
CTG GCG GCG GCG GTC CCC ACT GCT CCT GCT CCT TCC CCG ACG GTC ACT TGG ACT CCG GCG 205

E   P   G   P   A   L   N   Y   P   Q   E   E   A   T   L   N   E   M   F   R   52
GAG CCG GGC CCA GCT CTC AAC TAC CCT CAG GAG GAA GCT ACG CTC AAT GAG ATG TTT CGA 265

E   V   E   E   L   M   E   D   T   Q   H   K   L   R   S   A   V   E   E   M   72
GAG GTG GAG GAG CTG ATG GAA GAC ACT CAG CAC AAA CTG CGC AGT GCC GTG GAG GAG ATG 325

E   A   E   E   A   A   A   K   T   S   S   E   V   N   L   A   S   L   P   P   92
GAG GCG GAA GAA GCA GCT GCT AAA ACG TCC TCT GAG GTG AAC CTG GCA AGC TTA CCT CCC 385

N   Y   H   N   E   T   S   T   E   T   R   V   G   N   N   T   V   H   V   H   112
AAC TAT CAC AAT GAG ACC AGC ACG GAG ACC AGG GTG GGA AAT AAC ACA GTC CAT GTG CAC 445

Q   E   V   H   K   I   T   N   N   Q   S   G   Q   V   V   F   S   E   T   V   132
CAG GAA GTT CAC AAG ATA ACC AAC AAC CAG AGT GGA CAG GTG GTC TTT TCT GAG ACA GTC 505

I   T   S   V   G   D   E   E   G   K   R   S   H   E   C   I   I   D   E   D   152
ATT ACA TCT GTA GGG GAT GAA GAA GGC AAG AGG AGC CAT GAA TGT ATC ATT GAT GAA GAC 565

C   G   P   T   R   Y   C   Q   F   S   S   F   K   Y   T   C   Q   P   C   R   172
TGT GGG CCC ACC AGG TAC TGC CAG TTC TCC AGC TTC AAG TAC ACC TGC CAG CCA TGC CGG 625
  
```


FIG. 8B

D	Q	Q	M	L	C	T	R	D	S	E	C	C	G	D	Q	L	C	A	W	192
GAC	CAG	CAG	ATG	CTA	TGC	ACC	CGA	GAC	AGT	GAG	TGC	TGT	GGA	GAC	CAG	CTG	TGT	GCC	TGG	685
G	H	C	T	Q	K	A	T	K	G	G	N	G	T	I	C	D	N	Q	R	212
GGT	CAC	TGC	ACC	CAA	AAG	GCC	ACC	AAA	GGT	GGC	AAT	GGG	ACC	ATC	TGT	GAC	AAC	CAG	AGG	745
D	C	Q	P	G	L	C	C	A	F	Q	R	G	L	L	F	P	V	C	T	232
GAT	TGC	CAG	CCT	GGC	CTG	TGT	TGT	GCC	TTC	CAA	AGA	GGC	CTG	CTG	TTC	CCC	GTG	TGC	ACA	805
P	L	P	V	E	G	E	L	C	H	D	P	T	S	Q	L	L	D	L	I	252
CCC	CTG	CCC	GTG	GAG	GGA	GAG	CTC	TGC	CAT	GAC	CCC	ACC	AGC	CAG	CTG	CTG	GAT	CTC	ATC	865
T	W	E	L	E	P	E	G	A	L	D	R	C	P	C	A	S	G	L	L	272
ACC	TGG	GAA	CTG	GAG	CCT	GAA	GGA	GCT	TTG	GAC	CGA	TGC	CCC	TGC	GCC	AGT	GGC	CTC	CTA	925
C	Q	P	H	S	H	S	L	V	Y	M	C	K	P	A	F	V	G	S	H	292
TGC	CAG	CCA	CAC	AGC	CAC	AGT	CTG	GTG	TAC	ATG	TGC	AAG	CCA	GCC	TTC	GTG	GGC	AGC	CAT	985
D	H	S	E	E	S	Q	L	P	R	E	A	P	D	E	Y	E	D	V	G	312
GAC	CAC	AGT	GAG	GAG	AGC	CAG	CTG	CCC	AGG	GAG	GCC	CCG	GAT	GAG	TAC	GAA	GAT	GTT	GGC	1045
F	I	G	E	V	R	Q	E	L	E	D	L	E	R	S	L	A	Q	E	M	332
TTC	ATA	GGG	GAA	GTG	CGC	CAG	GAG	CTG	GAA	GAC	CTG	GAG	CGG	AGC	CTA	GCC	CAG	GAG	ATG	1105

FIG. 8C

A F E G P A P V E S L G G E E I * 350
GCA TTT GAG GGG CCT GCC CCT GTG GAG TCA CTA GGC GGA GAG GAG ATT TAG 1159
GCCCAGACCCAGCTGAGTCACTGGTAGATGTGCAATAGAAATGGCTAATTATTTTCCCAGGAGTGTCCTCCCAAGTGTTGG 1238
AATGGCCCGCAGCTCCTTCCCAGTAGCTTTTCCCTCTGGCTTGACAAGGTACAGTGCAGTACATTCTTCCAGCCGCCCTG 1317
CTTCTCTGACTTGGGAAAGACAGGCATGGCGGTAAGGCGAGCGGTGAGTCGTCCTCGCTGTGCTAGAAACGCTGTC 1396
TTGTTCTTCATGGATGGAAAGATTGTGTTGAAGGAGAGGATGGGAAGGGGTGAAGTCTGCTCATGATGGATTGTGGGGGA 1475
TACAGGGAGGAGGATGCCCTTGCCCTTGCCAGACGTGGACTTGGCAAAATGTAAACCTTTGCTTTGTCTGTGCGCGCTCCCAT 1554
GGGCTGAGGCAGTGGCTACACAAGAGCTATGCTGCTGTGGCCTCCACATATTTCATCCCTGTGTTTCAGCTCCTACC 1633
TCACTGTCAGCACAGCCCTTCATAGCCACGCCCTCTGCTCACCACAGCCCTAGGAGGGGACCAGAGGGGACTTCTCT 1712
CAGAGCCCCATGCTCTCTCAACCCCCATACCCAGCCTCTGTGCCAGCGACAGTCCCTTCCAAATGGAGGGAGTGAAAT 1791
CCTTTGGTTTAAATTATTTTCTCCTTCAAGGCACGCCCTGCCACTAAGGTCAGGCTGACTTGCAATGTCCCTCTAACGTTTCG 1870
TAGCAGTGTGGTGGACACTGTCTTCCACCGACTGCTTCAATACCTCTGAAGCCAGTGTCTGGAGTGCAGTTCGTGTAA 1949
ATTAATTGCAGGAAGTACTTGGCTAATTGTAGGGCTAGGATTGTGAATGAAATTTCGAAAGTCGCTTAGCAACAAT 2028
GGAAAGCCTTTCTCAGTCACACCGAGAAGTCACAACCAAGCCAGGTTGTGTAGAGTACAGCTGTGACATACAGACAGAA 2107

FIG. 8D

GAAGGCTGGGCTGGATGTCAGGCCCTCAGATGACGGTTTCAGGTGCCAGGAACCTATTACCAATTCGTATCTATCCAGAGT 2186
TATTAAAAATTGAAAGTTGCACACATTTGTATAAGCATGCCCTTCTCCTGAGTTTAAATATATATGATACACAAAACATG 2265
TGGCCCTCAAAGATCATGCACAAAACCACTACTCTTTTGCTAATTCTTGGACTTTTCTCTTTGATTTTCAATAAATACAAA 2344
TCCCCTTCATGCAAAAAAATAAGGGGGCGCGCGC 2381